- 5'-CCATCCTCTTGAAAATCTC-3' ① (SEQ ID NO:5)
- 5'-TCTCRTCTCACAAGTTTGGC-3' @ (SEQ ID NO:6).
- (6) The sixth invention is an oligonucleotide characterized in that the gene sequence of a spacer region between a gene coding 16S rRNA and a gene coding 23S rRNA of Pectinatus cerevisiiphilus has at least one of the following sequence group or the corresponding complementary sequence:
 - 5'-CACTCTTACAAGTATCTAC-3' ③ (SEQ ID NO:7)
 - 5'-CCACAATATTTCCGACCAGC-3' 4 (SEQ ID NO:8)
 - 5'-AGTCTTCTCTACTGCCATGC-3' (SEQ ID NO:9).--

Please replace the two paragraphs beginning at page 4, line 15, and ending at page 4, line 22, with the following rewritten paragraphs:

- --(11) The eleventh invention is a method as in (9), wherein the nucleotide sequence coding the 16S rRNA gene of Pectinatus frisingensis has the following sequence:
 - 5'-CGTATCCAGAGATGGATATT-3' © (SEQ ID NO:10)
- (12) The twelfth invention is a method as in (10), wherein the nucleotide sequence coding the 16S rRNA gene of Pectinatus cerevisiiphilus has the following sequence:

IN THE CLAIMS

Please amend claims 5, 6, 11 and 12 as follows: